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43		44		46		48		50		52		1		2	
UniGene		Clone		Name		BNS_Sym		BNS_Desc		BNS_Chrom		TC..U...		UA...	
R1	1	Invasive Ability												4.9	
	2	Vasculogenic Mimicry												122.0	
	3	Cell Motility													
	4	Sex													
R2	5	Hs.23590		solute carrier f		SLC16A4		solute carrier f		1p12					
	6	Hs.93574		homeo box D3		HOXD3		homeo box D3		2q31-q37					
	7	Hs.662		cerebellin 1 pr		CB									
	8	Hs.89414		chemokine (C-		CX									
R3	9	Hs.91585		multispanning		TM									
	10	Hs.77613		ataxia telangie		AT									
	11	Hs.75721		profilin 1		PFN1		profilin 1		17p13.3					
	12	Hs.2714		forkhead (Dros		FOXG1B		forkhead BOX		14q12-q13					
R4	13	Hs.247469		ESTs, Highly s											
	27	Hs.2384		tumor protein		TPD52		tumor protein		8q21					
	29	Hs.183656		Ests		VNN3		vanin 3		6q23-q24					
	31	Hs.14355		Ests											
R5	32	Hs.15740		Homo sapiens											
	33	Hs.182817		Ests											
	34	Hs.241065		296											
	35	Hs.82572		281											
R6	36	Hs.78465		358											
	37	Hs.22466		809											
	38	Hs.155101		502											
	59	Hs.84746		724615.0		chromosome c		CHC1		chromosome c		1p36.1			

FIG. 2

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UniGene	Clone	Name	BNS_Symb	BNS_Desc	BNS_Ch	TC..	U...	UA...	U...	18g1	20r1	20	U...	A...	U...	U...	M9...	M9...
1	Invasive Ability							4.9	4.2	U...								2.6
2	Vasulogenic Mimicry								+									-
3	Cell Motility								122.0									
4	Sex																	
5	Hs.79033	H.sapiens mR	QPC	glutaminyl-pep	2p22.3													
6	Hs.216726	ESTs																
7	Hs.34578	alpha2,3-sialyl	ST3GALM	alpha2,3-sialyl	3q12.2													
8	Hs.154069	melan-A	MILAN	melan-A	9p24.2													
9	Hs.23202	ESTs																
10	Hs.74561	alpha-2-macro	A2M	alpha-2-macro	12p13.3-p1													
11	Hs.83384	S100 calcium	S100B	S100 calcium	21q22.3													
12	Hs.8180	syndecan bindi	SDCBP	syndecan bindi	8q12													
13	Hs.74561	alpha-2-macro	A2M	alpha-2-macro	12p13.3-p1													
27	Hs.1975	ESTs, Weakly	FLJ21007	hypothetical pr	13q21.1													
28	Hs.107573	sialyltransferase																
29	Hs.29189	ESTs	ATP11A	ATPase, Class	13q34													
30	Hs.1940	crystallin, alph	CRYAB	crystallin, alph	11q22.3-q													
31	Hs.17752	ESTs, Highly s	PS-PLA1	phosphatidylse	3q13.13-q1													
32	Hs.69547	myelin basic p	MBP	myelin basic p	18q23													
33	Hs.78361	glycoprotein M																
34	Hs.241532	synixin 7																
35	Hs.1244	CD9 antigen (p	CD9	CD9 antigen (p	12p13													
36	Hs.170133	forkhead (Dros	FOXO1A	forkhead box	13q14.1													
37	Hs.82845	Human clone 2																
38	Hs.240615	CD68 antigen	NDRG3	NDRG family	20q11.21-q													
59	Hs.239459	regulatory solu	RSC1A1	regulatory solu	1p36.1													

Sorted on this column,  
 Sorting ONLY the rows  
 containing expression data

FIG. 3

FIG. 4

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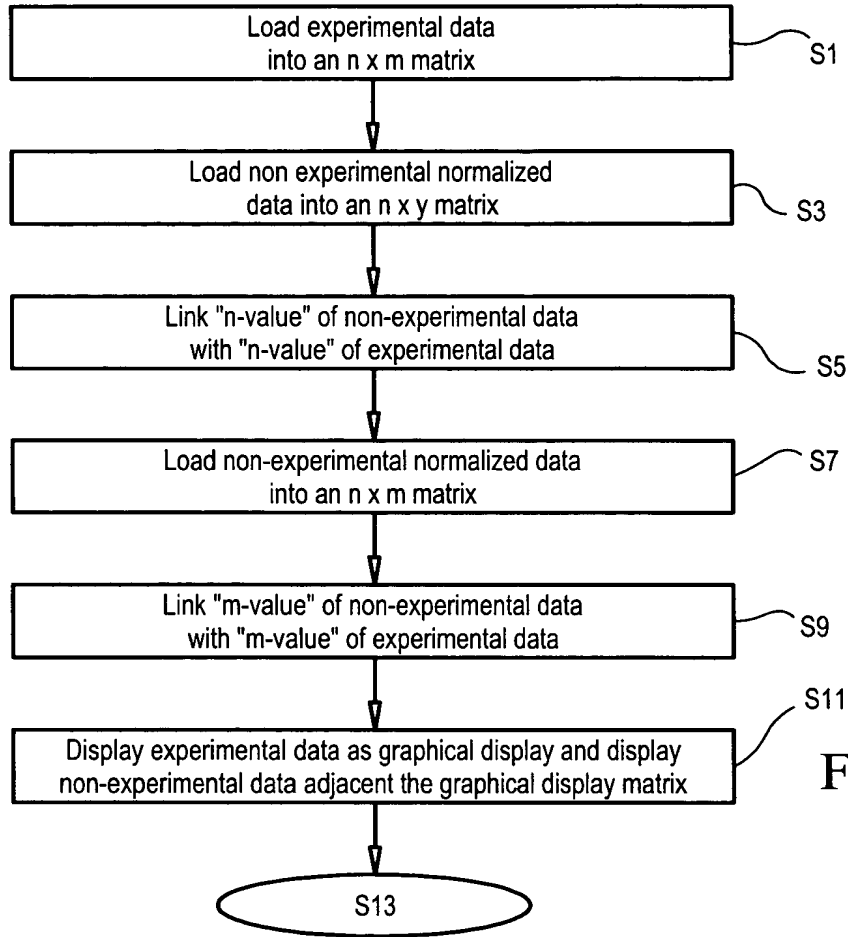


FIG. 5A

	101	102	103
201	101,201	102,201	103,201
202	101,202	102,202	103,202
203	101,203	102,203	103,203
204	101,204	102,204	103,204

FIG. 6A

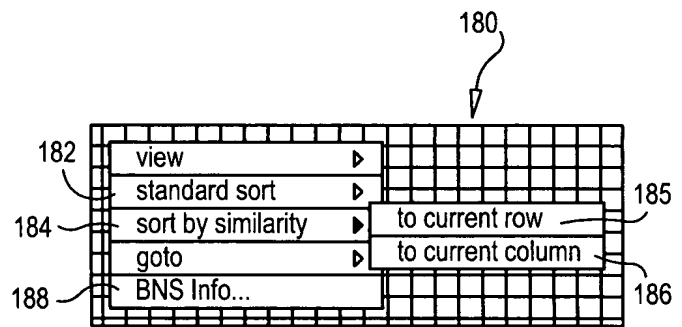
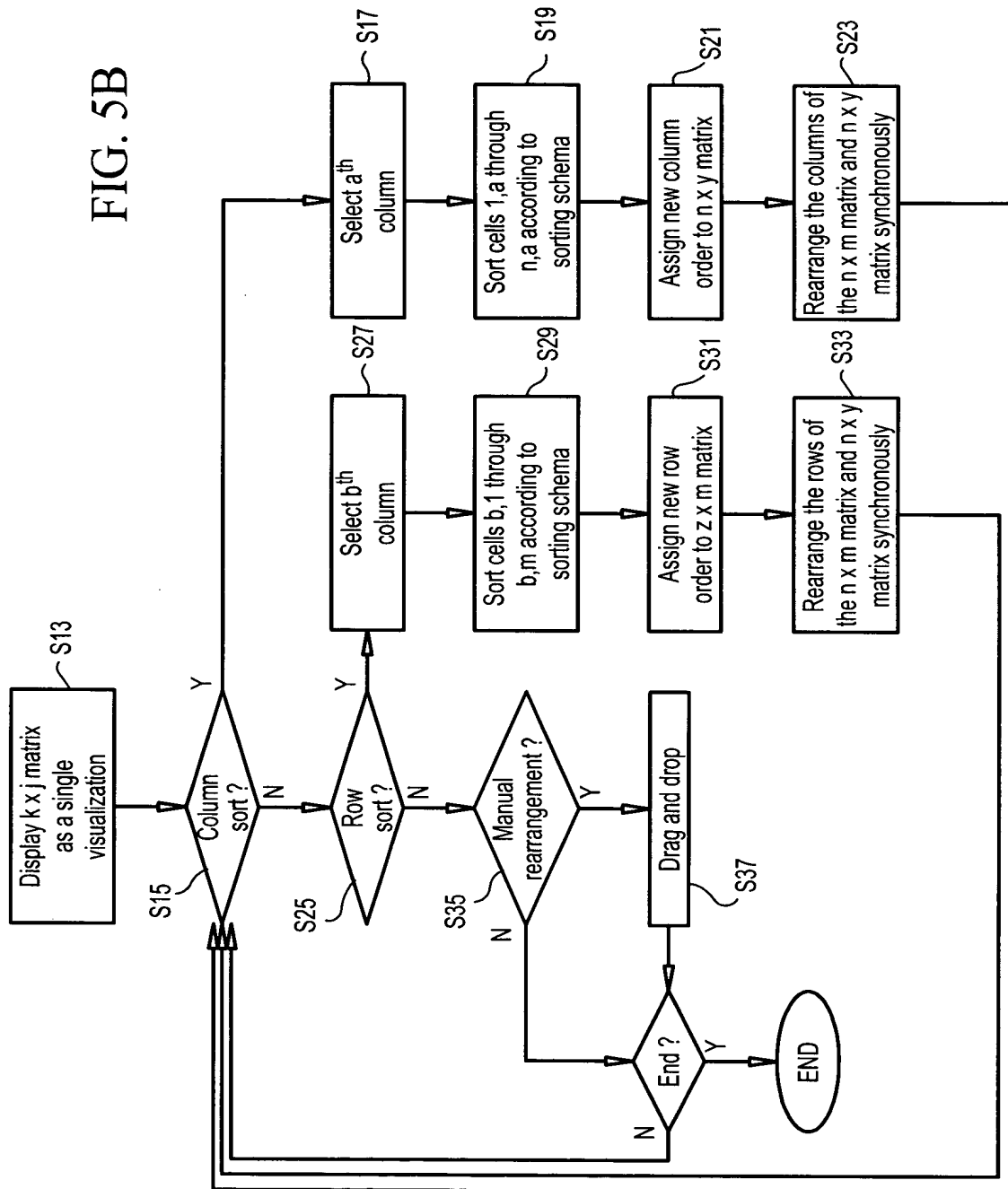


FIG. 6B

FIG. 5B



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	101	102	103
202	101,202	102,202	103,202
201	101,201	102,201	103,201
203	101,203	102,203	103,203
204	101,203	102,203	103,203

FIG. 6C

	101	102	103
202	101,202	102,202	103,202
203	101,203	102,203	103,203
201	101,201	102,201	103,201
204	101,204	102,204	103,204

FIG. 6D

Sort Settings

Positive Value

8.0

302

(fold increase)

300

Negative Value

8.0

304

(fold decrease)

Window Size

2

306

(num columns)

Increment

1

308

(num columns)

Similarity Increment

100

310

(num rows)

OK

Cancel

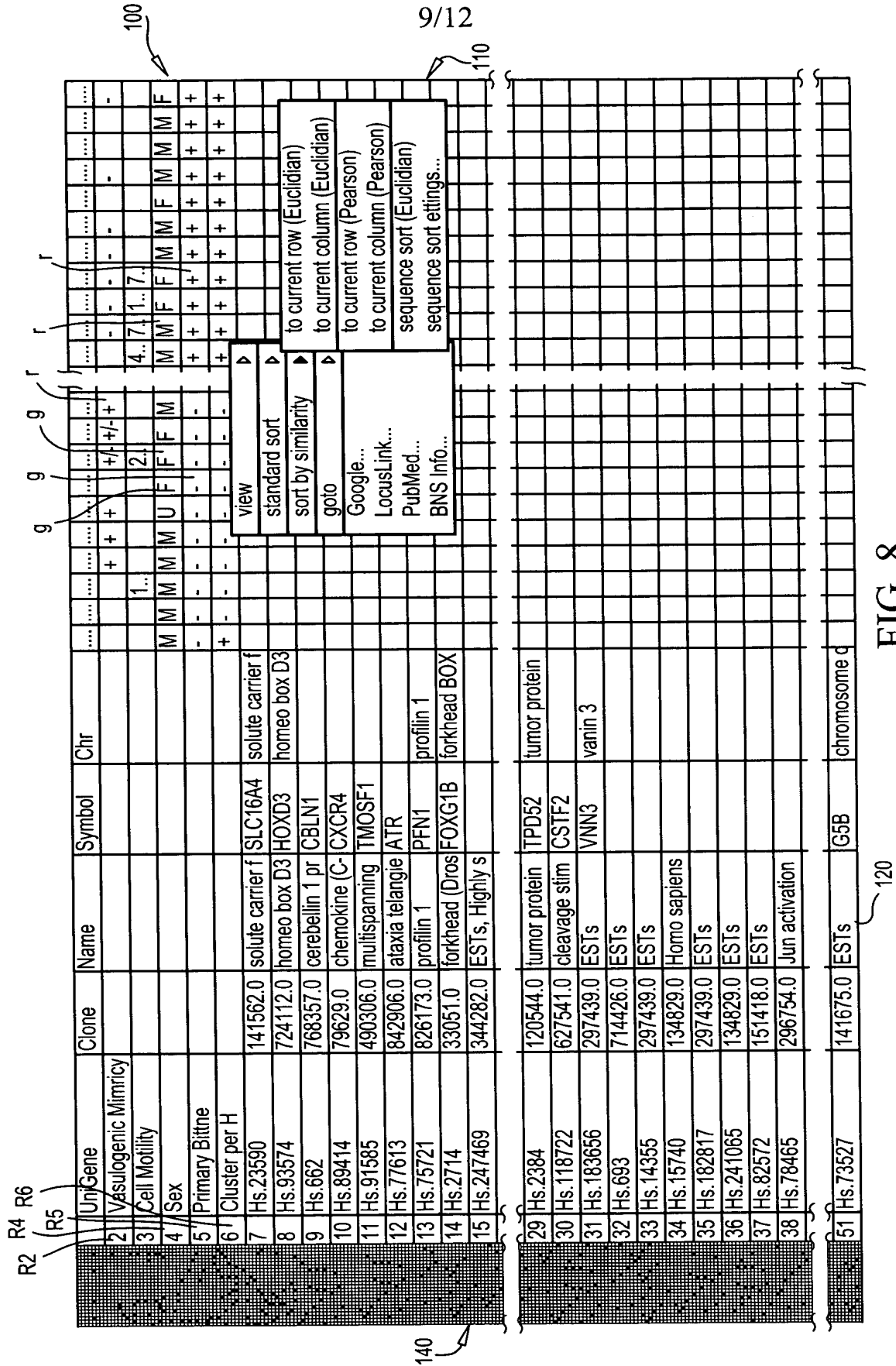
FIG. 9A

+	+	-		+	-	-	+	+	+	320
8.0	8.0	1/8	1	8.0	1/8	1/8	8.0	8.0	8.0	322
2.1	0.3	0.1	0.2	2.3	0.4	0.1	1.1	0.7	2.2	324
0.3	2.4	0.6	0.1	1.3	0.2	0.2	0.4	1.2	1.0	326
0.3	1.2	0.4	0.1	1.1	0.5	0.2	0.6	1.1	0.8	328

FIG. 9B







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UniGene	Clone	Name	Symbol	Chr	g										r										100										110																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																												
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FIG. 10

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C1	C2	C3	C4	C5	C6	C7	C8	C9	C10	C11	C12	C13	C14

FIG. 11A

C1	C2	C3	C4	C5	C6	C7	C8	C9	C10	C11	C12	C13	C14
1	1	1	1	8	8	8	8	1	1	1	1	1	1

← 332

FIG. 11B

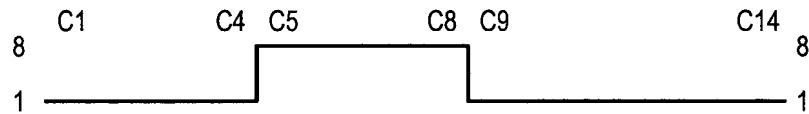


FIG. 11C

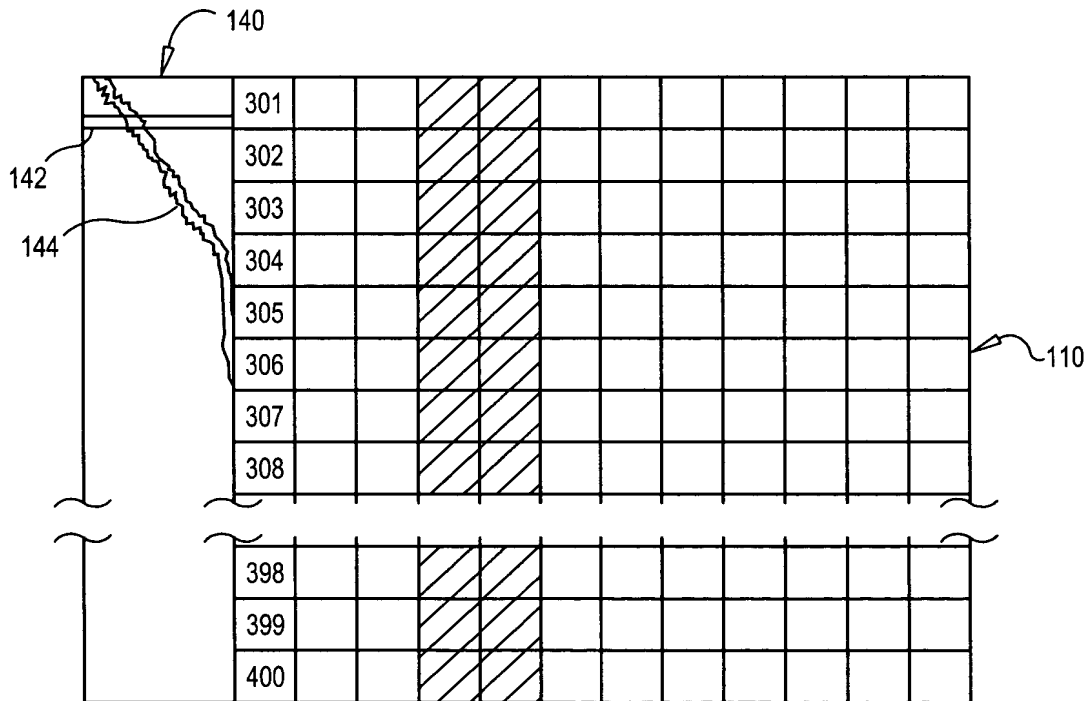


FIG. 13

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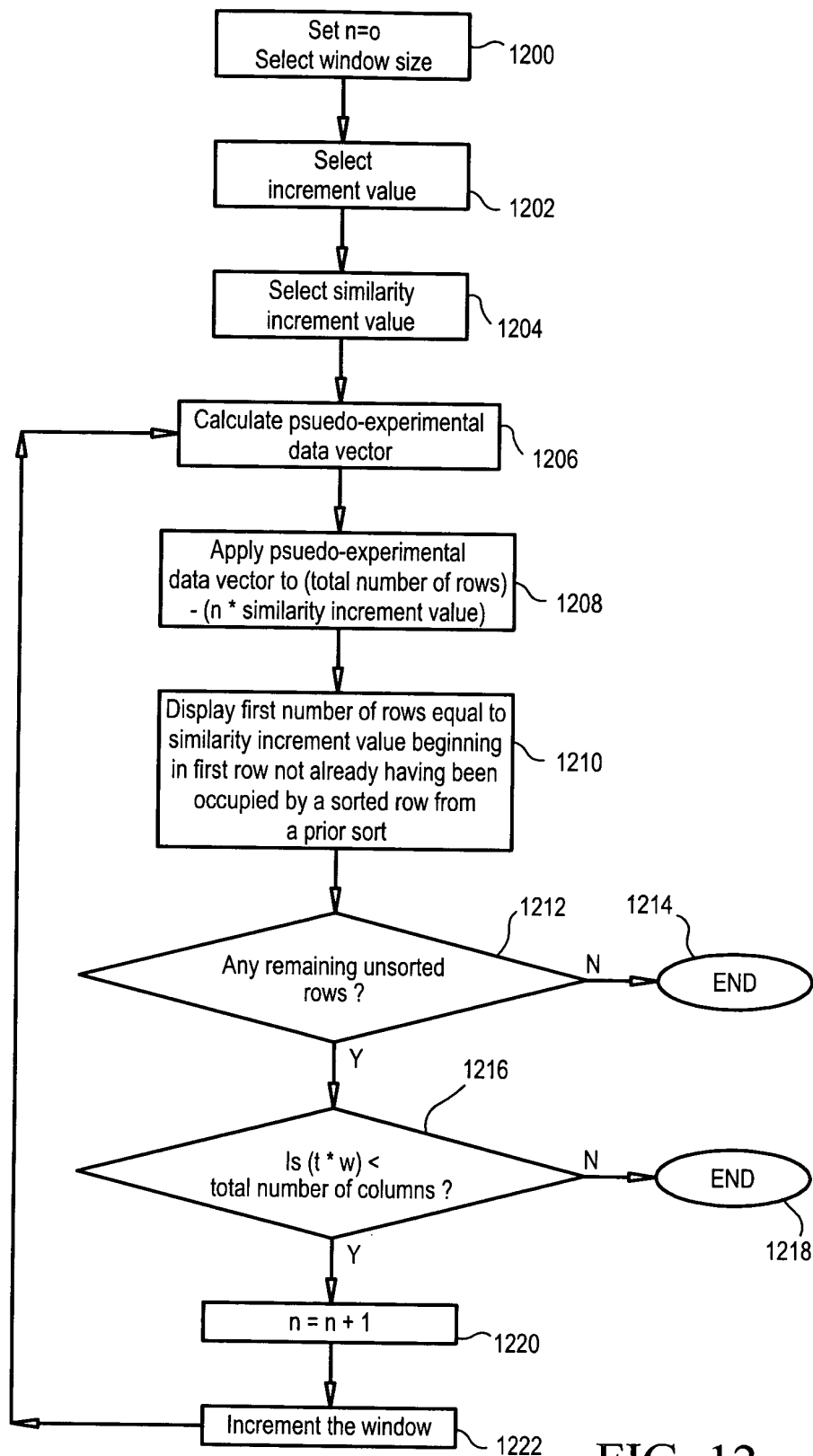


FIG. 12